

us-09-090-672b-1.oligo\_1.rst

Fri Apr 7 09:12:41 2000

GenCore version 4.5  
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OM nucleic - nucleic search, using sw model  
Run on: April 5, 2000, 18:38:42 ; Search time 6423.57 Seconds  
(without alignments)  
2513.359 Million cell updates/sec

Title: US-09-090-672B-1  
Perfect score: 4276  
Sequence: 1 TTCACGCTTTTTCCTGC.....ATCAGAAAAA 4276

Scoring table: OLIGO\_NUC  
Gapop 60.0 , Gapext 60.0

Searched: 4538634 seqs, 1887831982 residues

Word size : 0  
Total number of hits satisfying chosen parameters: 9077268

Minimum DB seq length: 0  
Maximum DB seq length: 1000000  
Post-processing: Listing first 45 summaries

Database : EST:  
1: em\_est1:  
2: em\_est2:  
3: em\_est3:  
4: em\_est4:  
5: em\_est5:  
6: em\_est6:  
7: em\_est7:  
8: em\_est8:  
9: em\_est9:  
10: em\_est10:  
11: em\_est11:  
12: em\_est12:  
13: em\_est13:  
14: em\_est14:  
15: em\_est15:  
16: em\_est16:  
17: em\_est17:  
18: em\_est18:  
19: em\_est19:  
20: gb\_est1:  
21: gb\_est2:  
22: gb\_est3:  
23: gb\_est4:  
24: gb\_est5:  
25: gb\_est6:  
26: gb\_est7:  
27: gb\_est8:  
28: gb\_est9:  
29: gb\_est10:  
30: gb\_est11:  
31: gb\_est12:  
32: gb\_est13:  
33: gb\_est14:  
34: gb\_est15:  
35: gb\_est16:  
36: gb\_est17:  
37: gb\_est18:  
38: gb\_est19:  
39: gb\_est20:  
40: gb\_est21:  
41: gb\_est22:  
42: gb\_est23:  
43: gb\_est24:  
44: gb\_est25:  
45: gb\_est26:  
46: gb\_est27:  
47: gb\_est28:  
48: gb\_est29:  
49: gb\_est30:  
50: gb\_est31:  
51: gb\_est32:  
52: em\_est20:  
53: em\_est21:  
54: em\_est22:  
55: em\_est23:  
56: em\_est24:  
57: em\_est25:  
58: em\_est26:  
59: gb\_est33:  
60: gb\_est34:  
61: gb\_est35:  
62: gb\_est36:  
63: gb\_est37:  
64: gb\_est38:  
65: em\_est27:  
66: em\_est28:  
67: em\_est29:  
68: em\_est30:  
69: gb\_est39:  
70: gb\_est40:  
71: gb\_est41:  
72: gb\_est42:  
73: gb\_est43:  
74: gb\_est44:  
75: em\_est31:  
76: em\_est32:  
77: em\_est33:  
78: em\_est34:  
79: gb\_gss1:  
80: gb\_gss2:  
81: gb\_gss3:  
82: gb\_gss4:  
83: em\_gss1:  
84: em\_gss2:  
85: em\_gss3:  
86: em\_gss4:  
87: gb\_gss5:  
88: gb\_gss6:  
89: gb\_gss7:  
90: gb\_gss8:  
91: gb\_gss9:  
92: em\_gss5:  
93: em\_gss6:  
94: em\_gss7:  
95: em\_gss8:  
96: em\_gss9:  
97: em\_gss10:  
98: em\_gss11:  
99: gb\_gss10:  
100: gb\_gss11:  
101: em\_gss12:  
102: gb\_gss12:  
103: gb\_gss13:  
104: gb\_gss14:  
105: gb\_gss15:

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | ID | Description        |
|------------|-------|-------------|--------|----|--------------------|
| C 1        | 386   | 9.0         | 506    | 45 | AI392674           |
|            |       |             |        |    | AI392674 t947c02.x |